RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: _10/650,326β_

Source: _1FWO_

Date Processed by STIC: _8/20/05_

ENTERED

IFWO

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 3 <110> APPLICANT: Hruska, Keith A.
 4        McCartney, John E.
 5        Charette, Marc F.
 7 <120> TITLE OF INVENTION: CONJOINT ADMINISTRATION OF MORPHOGENS AND ACE INHIBITORS IN
 8        TREATMENT OF CHRONIC RENAL FAILURE
10 <130> FILE REFERENCE: JJJ-P01-599
12 <140> CURRENT APPLICATION NUMBER: 10/650,326B
13 <141> CURRENT FILING DATE: 2003-08-28
15 <150> PRIOR APPLICATION NUMBER: 60/406,431
16 <151> PRIOR FILING DATE: 2002-08-28
18 <160> NUMBER OF SEQ ID NOS: 31
20 <170> SOFTWARE: PatentIn version 3.2
22 <210> SEQ ID NO: 1
23 <211> LENGTH: 139
24 <212> TYPE: PRT
25 <213> ORGANISM: Homo sapiens
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30 1               5                   10                  15
33 Asn Gln Glu Ala Leu Arg Met Ala Asn Val Ala Glu Asn Ser Ser Ser
34              20                  25                  30
37 Asp Gln Arg Gln Ala Cys Lys Lys His Glu Leu Tyr Val Ser Phe Arg
38          35                  40                  45
41 Asp Leu Gly Trp Gln Asp Trp Ile Ile Ala Pro Glu Gly Tyr Ala Ala
42      50                  55                  60
45 Tyr Tyr Cys Glu Gly Glu Cys Ala Phe Pro Leu Asn Ser Tyr Met Asn
46 65                  70                  75                  80
49 Ala Thr Asn His Ala Ile Val Gln Thr Leu Val His Phe Ile Asn Pro
50              85                  90                  95
53 Glu Thr Val Pro Lys Pro Cys Cys Ala Pro Thr Gln Leu Asn Ala Ile
54              100                 105                 110
57 Ser Val Leu Tyr Phe Asp Asp Ser Ser Asn Val Ile Leu Lys Lys Tyr
58          115                 120                 125
61 Arg Asn Met Val Val Arg Ala Cys Gly Cys His
62      130                 135
65 <210> SEQ ID NO: 2
66 <211> LENGTH: 97
67 <212> TYPE: PRT
68 <213> ORGANISM: Homo sapiens
70 <400> SEQUENCE: 2
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76 Leu Ser Ile Leu Gly Leu Pro His Arg Pro Arg Pro His Leu Gln Gly
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77                    20                        25                        30
80 Lys His Asn Ser Ala Pro Met Phe Met Leu Asp Leu Tyr Asn Ala Met
81              35                    40                        45
84 Ala Val Glu Glu Gly Gly Gly Pro Gly Gly Gln Gly Phe Ser Tyr Pro
85       50                        55                    60
88 Tyr Lys Ala Val Phe Ser Thr Gln Gly Pro Pro Leu Ala Ser Leu Gln
89 65                    70                        75                    80
92 Asp Ser His Phe Leu Thr Asp Ala Asp Met Val Met Ser Phe Val Asn
93                    85                        90                    95
96 Leu
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100 <211> LENGTH: 431
101 <212> TYPE: PRT
102 <213> ORGANISM: Homo sapiens
104 <400> SEQUENCE: 3
106 Met His Val Arg Ser Leu Arg Ala Ala Ala Pro His Ser Phe Val Ala
107 1               5                        10                        15
110 Leu Trp Ala Pro Leu Phe Leu Leu Arg Ser Ala Leu Ala Asp Phe Ser
111              20                        25                        30
114 Leu Asp Asn Glu Val His Ser Ser Phe Ile His Arg Arg Leu Arg Ser
115              35                        40                        45
118 Gln Glu Arg Arg Glu Met Gln Arg Glu Ile Leu Ser Ile Leu Gly Leu
119       50                        55                        60
122 Pro His Arg Pro Arg Pro His Leu Gln Gly Lys His Asn Ser Ala Pro
123 65                    70                        75                    80
126 Met Phe Met Leu Asp Leu Tyr Asn Ala Met Ala Val Glu Glu Gly Gly
127                    85                        90                    95
130 Gly Pro Gly Gly Gln Gly Phe Ser Tyr Pro Tyr Lys Ala Val Phe Ser
131              100                       105                       110
134 Thr Gln Gly Pro Pro Leu Ala Ser Leu Gln Asp Ser His Phe Leu Thr
135              115                       120                       125
138 Asp Ala Asp Met Val Met Ser Phe Val Asn Leu Val Glu His Asp Lys
139       130                       135                       140
142 Glu Phe Phe His Pro Arg Tyr His His Arg Glu Phe Arg Phe Asp Leu
143 145                       150                       155                       160
146 Ser Lys Ile Pro Glu Gly Glu Ala Val Thr Ala Ala Glu Phe Arg Ile
147              165                       170                       175
150 Tyr Lys Asp Tyr Ile Arg Glu Arg Phe Asp Asn Glu Thr Phe Arg Ile
151              180                       185                       190
154 Ser Val Tyr Gln Val Leu Gln Glu His Leu Gly Arg Glu Ser Asp Leu
155              195                       200                       205
158 Phe Leu Leu Asp Ser Arg Thr Leu Trp Ala Ser Glu Glu Gly Trp Leu
159       210                       215                       220
162 Val Phe Asp Ile Thr Ala Thr Ser Asn His Trp Val Val Asn Pro Arg
163 225                       230                       235                       240
166 His Asn Leu Gly Leu Gln Leu Ser Val Glu Thr Leu Asp Gly Gln Ser
167              245                       250                       255
170 Ile Asn Pro Lys Leu Ala Gly Leu Ile Gly Arg His Gly Pro Gln Asn
171              260                       265                       270
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174 Lys Gln Pro Phe Met Val Ala Phe Phe Lys Ala Thr Glu Val His Phe
175         275                 280                 285
178 Arg Ser Ile Arg Ser Thr Gly Ser Lys Gln Arg Ser Gln Asn Arg Ser
179     290                 295                 300
182 Lys Thr Pro Lys Asn Gln Glu Ala Leu Arg Met Ala Asn Val Ala Glu
183 305                 310                 315                 320
186 Asn Ser Ser Ser Asp Gln Arg Gln Ala Cys Lys Lys His Glu Leu Tyr
187                 325                 330                 335
190 Val Ser Phe Arg Asp Leu Gly Trp Gln Asp Trp Ile Ile Ala Pro Glu
191             340                 345                 350
194 Gly Tyr Ala Ala Tyr Tyr Cys Glu Gly Glu Cys Ala Phe Pro Leu Asn
195         355                 360                 365
198 Ser Tyr Met Asn Ala Thr Asn His Ala Ile Val Gln Thr Leu Val His
199     370                 375                 380
202 Phe Ile Asn Pro Glu Thr Val Pro Lys Pro Cys Cys Ala Pro Thr Gln
203 385                 390                 395                 400
206 Leu Asn Ala Ile Ser Val Leu Tyr Phe Asp Asp Ser Ser Asn Val Ile
207             405                 410                 415
210 Leu Lys Lys Tyr Arg Asn Met Val Val Arg Ala Cys Gly Cys His
211             420                 425                 430
214 <210> SEQ ID NO: 4
215 <211> LENGTH: 139
216 <212> TYPE: PRT
217 <213> ORGANISM: Mus musculus
219 <400> SEQUENCE: 4
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225 Asn Gln Glu Ala Leu Arg Met Ala Ser Val Ala Glu Asn Ser Ser Ser
226             20                  25                  30
229 Asp Gln Arg Gln Ala Cys Lys Lys His Glu Leu Tyr Val Ser Phe Arg
230         35                  40                  45
233 Asp Leu Gly Trp Gln Asp Trp Ile Ile Ala Pro Glu Gly Tyr Ala Ala
234     50                  55                  60
237 Tyr Tyr Cys Glu Gly Glu Cys Ala Phe Pro Leu Asn Ser Tyr Met Asn
238 65              70                  75                  80
241 Ala Thr Asn His Ala Ile Val Gln Thr Leu Val His Phe Ile Asn Pro
242             85                  90                  95
245 Asp Thr Val Pro Lys Pro Cys Cys Ala Pro Thr Gln Leu Asn Ala Ile
246             100                 105                 110
249 Ser Val Leu Tyr Phe Asp Asp Ser Ser Asn Val Ile Leu Lys Lys Tyr
250         115                 120                 125
253 Arg Asn Met Val Val Arg Ala Cys Gly Cys His
254     130                 135
257 <210> SEQ ID NO: 5
258 <211> LENGTH: 139
259 <212> TYPE: PRT
260 <213> ORGANISM: Homo sapiens
262 <400> SEQUENCE: 5
264 Ala Val Arg Pro Leu Arg Arg Arg Gln Pro Lys Lys Ser Asn Glu Leu
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265 1                5               10              15
268 Pro Gln Ala Asn Arg Leu Pro Gly Ile Phe Asp Asp Val His Gly Ser
269              20              25              30
272 His Gly Arg Gln Val Cys Arg Arg His Glu Leu Tyr Val Ser Phe Gln
273         35              40              45
276 Asp Leu Gly Trp Leu Asp Trp Val Ile Ala Pro Gln Gly Tyr Ser Ala
277      50              55              60
280 Tyr Tyr Cys Glu Gly Glu Cys Ser Phe Pro Leu Asp Ser Cys Met Asn
281 65              70              75              80
284 Ala Thr Asn His Ala Ile Leu Gln Ser Leu Val His Leu Met Lys Pro
285              85              90              95
288 Asn Ala Val Pro Lys Ala Cys Cys Ala Pro Thr Lys Leu Ser Ala Thr
289         100             105             110
292 Ser Val Leu Tyr Tyr Asp Ser Ser Asn Asn Val Ile Leu Arg Lys His
293         115             120             125
296 Arg Asn Met Val Val Lys Ala Cys Gly Cys His
297      130             135

300 <210> SEQ ID NO: 6
301 <211> LENGTH: 139
302 <212> TYPE: PRT
303 <213> ORGANISM: Mus musculus
305 <400> SEQUENCE: 6
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308 1                5               10              15
311 Pro His Pro Asn Lys Leu Pro Gly Ile Phe Asp Asp Gly His Gly Ser
312              20              25              30
315 Arg Gly Arg Glu Val Cys Arg Arg His Glu Leu Tyr Val Ser Phe Arg
316         35              40              45
319 Asp Leu Gly Trp Leu Asp Trp Val Ile Ala Pro Gln Gly Tyr Ser Ala
320      50              55              60
323 Tyr Tyr Cys Glu Gly Glu Cys Ala Phe Pro Leu Asp Ser Cys Met Asn
324 65              70              75              80
327 Ala Thr Asn His Ala Ile Leu Gln Ser Leu Val His Leu Met Lys Pro
328              85              90              95
331 Asp Val Val Pro Lys Ala Cys Cys Ala Pro Thr Lys Leu Ser Ala Thr
332         100             105             110
335 Ser Val Leu Tyr Tyr Asp Ser Ser Asn Asn Val Ile Leu Arg Lys His
336         115             120             125
339 Arg Asn Met Val Val Lys Ala Cys Gly Cys His
340      130             135

343 <210> SEQ ID NO: 7
344 <211> LENGTH: 588
345 <212> TYPE: PRT
346 <213> ORGANISM: Drosophila melanogaster
348 <400> SEQUENCE: 7
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354 Ile Val Arg Val Ala Ser Thr Glu Asp Ile Ser Gln Arg Phe Ile Ala
355              20              25              30
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358 Ala Ile Ala Pro Val Ala Ala His Ile Pro Leu Ala Ser Ala Ser Gly
359         35                  40                  45
362 Ser Gly Ser Gly Arg Ser Gly Ser Arg Ser Gly Gly Ala Ser Thr Ser
363         50                  55                  60
366 Thr Ala Leu Ala Lys Ala Phe Asn Pro Phe Ser Glu Pro Ala Ser Phe
367 65                  70                  75                  80
370 Ser Asp Ser Asp Lys Ser His Arg Ser Lys Thr Asn Lys Lys Pro Ser
371                 85                  90                  95
374 Lys Ser Asp Ala Asn Arg Gln Phe Asn Glu Val His Lys Pro Arg Thr
375             100                 105                 110
378 Asp Gln Leu Glu Asn Ser Lys Asn Met Ser Lys Gln Leu Val Asn Lys
379         115                 120                 125
382 Pro Asn His Asn Lys Met Ala Val Lys Glu Gln Arg Ser His His Lys
383         130                 135                 140
386 Lys Ser His His His Arg Ser His Gln Pro Lys Gln Ala Ser Ala Ser
387 145                 150                 155                 160
390 Thr Glu Ser His Gln Ser Ser Ser Ile Glu Ser Ile Phe Val Glu Glu
391                 165                 170                 175
394 Pro Thr Leu Val Leu Asp Arg Glu Val Ala Ser Ile Asn Val Pro Ala
395             180                 185                 190
398 Asn Ala Lys Ala Ile Ile Ala Glu Gln Gly Pro Ser Thr Tyr Ser Lys
399         195                 200                 205
402 Glu Ala Leu Ile Lys Asp Lys Leu Lys Pro Asp Pro Ser Thr Leu Val
403         210                 215                 220
406 Glu Ile Glu Lys Ser Leu Leu Ser Leu Phe Asn Met Lys Arg Pro Pro
407 225                 230                 235                 240
410 Lys Ile Asp Arg Ser Lys Ile Ile Ile Pro Glu Pro Met Lys Lys Leu
411                 245                 250                 255
414 Tyr Ala Glu Ile Met Gly His Glu Leu Asp Ser Val Asn Ile Pro Lys
415             260                 265                 270
418 Pro Gly Leu Leu Thr Lys Ser Ala Asn Thr Val Arg Ser Phe Thr His
419         275                 280                 285
422 Lys Asp Ser Lys Ile Asp Asp Arg Phe Pro His His His Arg Phe Arg
423         290                 295                 300
426 Leu His Phe Asp Val Lys Ser Ile Pro Ala Asp Glu Lys Leu Lys Ala
427 305                 310                 315                 320
430 Ala Glu Leu Gln Leu Thr Arg Asp Ala Leu Ser Gln Gln Val Val Ala
431             325                 330                 335
434 Ser Arg Ser Ser Ala Asn Arg Thr Arg Tyr Gln Val Leu Val Tyr Asp
435         340                 345                 350
438 Ile Thr Arg Val Gly Val Arg Gly Gln Arg Glu Pro Ser Tyr Leu Leu
439         355                 360                 365
442 Leu Asp Thr Lys Thr Val Arg Leu Asn Ser Thr Asp Thr Val Ser Leu
443         370                 375                 380
446 Asp Val Gln Pro Ala Val Asp Arg Trp Leu Ala Ser Pro Gln Arg Asn
447 385                 390                 395                 400
450 Tyr Gly Leu Leu Val Glu Val Arg Thr Val Arg Ser Leu Lys Pro Ala
451             405                 410                 415
454 Pro His His His Val Arg Leu Arg Arg Ser Ala Asp Glu Ala His Glu
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RAW SEQUENCE LISTING ERROR SUMMARY   DATE: 08/20/2005
PATENT APPLICATION: US/10/650,326B   TIME: 11:47:46

Input Set : **A:\JJJ-P01-599.txt**
Output Set: **N:\CRF4\08202005\J650326B.raw**

Please Note:
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the
Sequence Listing to ensure that a corresponding explanation is presented in the <220>
to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:24; Xaa Pos. 2,3,4,6,7,8,11,12,13,14,15,16,18,19,20,21,23,26,28,30,31
Seq#:24; Xaa Pos. 33,34,35,36,37,38,39,40,44,45,46,47,48,49,50,51,52,53,54
Seq#:24; Xaa Pos. 55,56,57,58,59,60,63,65,66,67,68,69,70,71,72,74,75,76,77
Seq#:24; Xaa Pos. 78,79,80,82,84,85,86,87,88,90,92,93,95,97
Seq#:25; Xaa Pos. 2,3,4,5,7,8,9,11,12,13,16,17,18,19,20,21,23,24,25,26,28
Seq#:25; Xaa Pos. 31,33,35,36,38,39,40,41,42,43,44,45,49,50,51,52,53,54,55
Seq#:25; Xaa Pos. 56,57,58,59,60,61,62,63,64,65,68,70,71,72,73,74,75,76,77
Seq#:25; Xaa Pos. 79,80,81,82,83,84,85,87,89,90,91,92,93,95,97,98,100,102
Seq#:26; Xaa Pos. 2,3,4,5
Seq#:27; Xaa Pos. 1,2,3,4,5,6,7,8,9,10,11,12,13,14,15,16,18,19,20,21,22,23
Seq#:27; Xaa Pos. 24,26,28,30,31,32,33,34,35,36,37,38,39,40,41,42,43,44,45
Seq#:27; Xaa Pos. 46,47,48,49,50,51,52,53,54,55,56,57,58,59,60,61,63,65,66
Seq#:27; Xaa Pos. 67,68,69,70,71,72,74,75,76,77,78,79,80,81,82,83,84,85,86
Seq#:27; Xaa Pos. 87,88,89,90,91,92,93,95,97
Seq#:28; Xaa Pos. 2,3,4,5,6,7,8,9,10,11,12,13,14,15,16,17,18,19,20,21,23,24
Seq#:28; Xaa Pos. 25,26,27,28,29,31,33,35,36,37,38,39,40,41,42,43,44,45,46
Seq#:28; Xaa Pos. 47,48,49,50,51,52,53,54,55,56,57,58,59,60,61,62,63,64,65
Seq#:28; Xaa Pos. 66,68,70,71,72,73,74,75,76,77,79,80,81,82,83,84,85,86,87
Seq#:28; Xaa Pos. 88,89,90,91,92,93,94,95,96,97,98,100,102
Seq#:29; Xaa Pos. 2,3,11,16,19,23,26,35,39,41,50,52,56,57,58,60,61,65,71,73
Seq#:29; Xaa Pos. 75,80,82,84,89,96
Seq#:30; Xaa Pos. 2,3

Invalid Line Length:
The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:24; Line(s) 2080
Seq#:25; Line(s) 2484
Seq#:26; Line(s) 2890
Seq#:27; Line(s) 2924
Seq#:28; Line(s) 3403
Seq#:29; Line(s) 3893

VERIFICATION SUMMARY   DATE: 08/20/2005
PATENT APPLICATION: **US/10/650,326B**   TIME: 11:47:46

Input Set : **A:\JJJ-P01-599.txt**
Output Set: **N:\CRF4\08202005\J650326B.raw**

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L:2451 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24 after pos.:0
M:341 Repeated in SeqNo=24
L:2856 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25 after pos.:0
M:341 Repeated in SeqNo=25
L:2914 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26 after pos.:0
L:3369 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 after pos.:0
M:341 Repeated in SeqNo=27
L:3405 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:3507 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:28
L:3741 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:28
L:3749 M:283 W: Missing Blank Line separator, <220> field identifier
L:3857 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:28
L:3859 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28 after pos.:0
M:341 Repeated in SeqNo=28
L:4039 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:29
L:4041 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:29 after pos.:0
M:341 Repeated in SeqNo=29
L:4083 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:30 after pos.:0
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